OIPE

RAW SEQUENCE LISTING

DATE: 09/06/2000

PATENT APPLICATION: US/09/645,706

TIME: 12:02:11

Input Set : A:\341-005US1.txt.txt

Output Set: N:\CRF3\09062000\I645706.raw

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ENTERED
See P. 5
      4 <110> APPLICANT: Wood, Keith V.
              Gruber, Monika G.
              Zhuang, Yao
      8 <120> TITLE OF INVENTION: SYNTHETIC NUCLEIC ACID MOLECULE
              COMPOSITIONS AND METHODS OF PREPARATION
     11 <130> FILE REFERENCE: 341.005US1
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/645,706
C--> 13 <141> CURRENT FILING DATE: 2000-08-24
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     19 <212> TYPE: DNA
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     25 ttagtagatg tgtttggtga cgaatcgctt tcctataaag agttttttga agctacatgc
                                                                              180
     26 ctcctagcgc aaagtctcca caattgtgga tacaagatga atgatgtagt gtcgatctgc
                                                                              240
                                                                              300
     27 gccgagaata ataaaagatt ttttattccc attattgcag cttggtatat tggtatgatt
     28 gtagcacctg ttaatgaaag ttacatccca gatgaactct gtaaggtcat gggtatatcg
     29 aaaccacaaa tagttttttg tacaaagaac attttaaata aggtattgga ggtacagagc
     30 agaactaatt toataaaaag gatoatoata ottgatactg tagaaaacat acacggttgt
     31 gaaagtette ecaatttat ttetegttat teggatggaa atattgeeaa etteaaacet
                                                                              540
     32 ttacattacg atcctgttga gcaagtggca gctatcttat gttcgtcagg cactactgga
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     33 ttaccgaaag gtgtaatgca aactcaccaa aatatttgtg tccgacttat acatgcttta
                                                                              660
     34 gaccccaggg caggaacgca acttattect ggtgtgacag tettagtata tetgeetttt
                                                                              720
     35 ttccatgctt ttgggttctc tataaacttg ggatacttca tggtgggtct tcgtgttatc
                                                                              780
                                                                              840
     36 atgttaagac gatttgatca agaagcattt ctaaaagcta ttcaggatta tgaagttcga
                                                                              900
     37 agtgtaatta acgttccagc aataatattg ttcttatcga aaagtccttt ggttgacaaa
                                                                              960
     1020
     39 qctgaqqttq caqtaaaacg attaaacttg ccaggaattc gctgtggatt tggtttgaca
                                                                             1080
     40 gaatctactt cagctaatat acacagtett ggggatgaat ttaaatcagg atcacttgga
     41 agagttactc ctttaatggc agctaaaata gcagataggg aaactggtaa agcattggga
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     42 ccaaatcaag ttggtgaatt atgcgttaaa ggtcccatgg tatcgaaagg ttacgtgaac
                                                                             1200
     43 aatgtagaag ctaccaaaga agctattgat gatgatggtt ggcttcactc tggagacttt
                                                                             1260
                                                                             1320
     44 ggatactatg atgaggatga gcatttctat gtggtggacc gttacaagga attgattaaa
                                                                                             SEP 20 2000
CIPE/JCWS
                                                                             1380
     45 tataagggct ctcaggtagc acctgcagaa ctagaagaga ttttattgaa aaatccatgt
                                                                             1440
     46 atcagagatg ttgctgtggt tggtattcct gatctagaag ctggagaact gccatctgcg
     47 tttgtggtta aacagcccgg aaaggagatt acagctaaag aagtgtacga ttatcttgcc
                                                                             1500
                                                                             1560
     48 gagagggtct cccatacaaa gtatttgcgt ggaggggttc gattcgttga tagcatacca
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     49 aggaatgtta caggtaaaat tacaagaaag gaacttctga agcagttgct ggagaagagt
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     53 <211> LENGTH: 1626
     54 <212> TYPE: DNA
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9/6/00

55 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING
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| 8 <223> OTHER INFORMATION: Sequence of clone YG#81-6G01  |            |  |  |  |  |  |  |
| 60 <400> SEQUENCE: 2   |            |  |  |  |  |  |  |
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| 02 Ctadeagety quyudutget offeeggee offeggaaaab accordates accordates   | .20        |  |  |  |  |  |  |
| os ceageagaeg eggeegge egaaeegee acceataaag agarrarya agarrarya -  | .80        |  |  |  |  |  |  |
| of coccayed adagected cadebyegga cacaagaaga aagaagaaga gargaaga  | 240        |  |  |  |  |  |  |
| os googagada acacaagade ecceaciono geracoguas ocosso   | 300        |  |  |  |  |  |  |
|  | 360        |  |  |  |  |  |  |
| or addocada cayeecedo guordayado nerrotario ayyoni yyon yyon   | 20         |  |  |  |  |  |  |
|  | 80         |  |  |  |  |  |  |
| os gadageocco condeceda coo objetat cossistant non-sistant non-  | 40         |  |  |  |  |  |  |
| 70 ccacacacag according a goad goad goad goad goad goad goad g   | 00         |  |  |  |  |  |  |
| 71 Claceguay gegenalgou autocomobile nations of the just the second  | 60         |  |  |  |  |  |  |
|  | 20         |  |  |  |  |  |  |
| /3 cccdatagett cegagette tataateett januareett ajjanjajan aanta et januareett  | 780        |  |  |  |  |  |  |
| 74 degeledge gallegaled agaagealed temmangeen evenjamen - Jamping  | 340        |  |  |  |  |  |  |
| 75 agegeaded aegecodes agenatuces construction and the construction of the constructio | 00         |  |  |  |  |  |  |
| 70 caogacteae caagecoaag ggaacegege vgoggogege   | 60         |  |  |  |  |  |  |
| 1/ googlaggoog cagoaaaacg accaaaccog coaggaacco googlaggoor - John - Joh | 20         |  |  |  |  |  |  |
| 70 gaacecace cagecaacae acadagecore agggangman an-maring marine and  | 080        |  |  |  |  |  |  |
| is agagecaece ecceaacygo agocaaaca yougasayyy aan oy o o o o o   | L40        |  |  |  |  |  |  |
| or coddaccdag coggegades acguateraa gystooning   | 200        |  |  |  |  |  |  |
| of date taged to accordance and the factor of the state o | 260        |  |  |  |  |  |  |
| or against and an arministration of the second seco | 320        |  |  |  |  |  |  |
|  | 380        |  |  |  |  |  |  |
| or accayagacy cogocycy cygoaccool galeetaynay cogyaynaan jaraan y  | 140        |  |  |  |  |  |  |
| os ceegeggeen andageoogg analysas asas estas amb sum   | 500        |  |  |  |  |  |  |
| ob gagagggee cecaeacaa geaecegege ggaggggee gaeeogaega engenemen   | 60         |  |  |  |  |  |  |
| or aggaregeta caggeratar ractargulary gallocooliga agoligotyto yyaningyty  | 20         |  |  |  |  |  |  |
| 00 334336  | 526        |  |  |  |  |  |  |
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| 93 <213> ORGANISM: Artificial Sequence   |            |  |  |  |  |  |  |
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| 96 <223> OTHER INFORMATION: Sequence of a synthetic luciferase   |            |  |  |  |  |  |  |
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| 100 ctgaccgccg gtgagatgtt gttccgtgct ctgcgtaaac attctcactt gcctcaagcc  | 120        |  |  |  |  |  |  |
| 101 ctggtggatg tcgtgggcga cgaaagcttg tcttataagg agtttttcga agctactgtc  | 180        |  |  |  |  |  |  |
| 102 ctgttggccc agtctctgca taattgcggt tacaaaatga acgatgtggt cagcatttgt  | 240        |  |  |  |  |  |  |
| 103 getgagaata acaccegett ttteateeca gtgattgeeg ettggtacat eggeatgatt  | 300        |  |  |  |  |  |  |
| 104 gtcgcccctg tgaatgaatc ttatatccca gacgagttgt gcaaggtcat gggtattagc  | 360        |  |  |  |  |  |  |
| 105 aaacetcaaa tegtgtttae taccaagaac attetgaata aagtettgga agtgeagtet  | 420        |  |  |  |  |  |  |
| 106 cgtactaact tcatcaagcg cattatcatt ctggataccg tcgagaatat ccacggctgt  | 480        |  |  |  |  |  |  |
| 107 gaaagettge caaactttat ttetegttat agegaeggta atategetaa etteaageet  | 540        |  |  |  |  |  |  |
| 108 etgeattttg atceagtgga geaagtegee getattttgt getetagegg cactaceggt  | 600<br>660 |  |  |  |  |  |  |
| 109 etgectaaag gegtgatgea gaeteaceaa aatatetgtg teegettgat teatgeeetg  | 000        |  |  |  |  |  |  |



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| 110 | gacccacgtg  | tgggtaccca   | gttgatccct  | ggcgtgactg   | tcctggtgta   | cttgccattc  | 720  |
|-----|-------------|--------------|-------------|--------------|--------------|-------------|------|
| 111 | tttcacgcct  | tcaatttttc   | tattaccctq  | ggctatttca   | tggtcggttt   | gcgcgtgatc  | 780  |
| 112 | atgtttcgtc  | gcttcgatca   | agaagetttt  | ctgaaggcca   | ttcaggacta   | cgaggtccgt  | 840  |
| 113 | agcgtgatca  | acqtcccttc   | tataatttta  | ttcctgagca   | aatctccatt   | ggtcgataag  | 900  |
| 114 | tatgacctga  | actetttaca   | cgaactgtgc  | tgtggcgctg   | cccctttggc   | taaagaggtg  | 960  |
| 115 | gccgaagtcg  | ctgccaagcg   | totgaatttg  | ccaggtatcc   | actacaactt   | tggtctgact  | 1020 |
| 116 | gagagcacct  | ctgctaacat   | tcatagettg  | cgtgatgaat   | tcaaatctqq   | cagcctgggt  | 1080 |
| 117 | cacataacte  | ctttgatggc   | cactaagato  | gccgaccgtg   | agaccggcaa   | agetetgggt  | 1140 |
| 118 | ccaaatcaag  | tcggcgaatt   | gtgtattaag  | ggtcctatgg   | totctaaagg   | ctacqtcaac  | 1200 |
| 119 | aatataaaaa  | ccactaagga   | agctatcgat  | gacgatggtt   | ggctgcacag   | cqqcqacttt  | 1260 |
| 120 | ggttattacg  | atgaggacga   | acatttctat  | gtcgtggatc   | gctacaaaga   | gttgattaag  | 1320 |
| 121 | tataaaggct  | ctcaggtcgc   | cccagctgag  | ctggaagaga   | tettgetgaa   | gaaccettge  | 1380 |
| 122 | attcatgaca  | taaccatcat   | gggtatccca  | gatttggaag   | ctggcgagct   | gcctagcgcc  | 1440 |
| 123 | tttatcataa  | aacaaccagg   | taaggaaatt  | accgctaaag   | aggtetacga   | ctatttggcc  | 1500 |
| 124 | gaacgcgtgt  | ctcacactaa   | atacctacat  | ggcggtgtcc   | acttcataga   | tagcatccct  | 1560 |
|     |             |              |             | gagttgctga   |              |             | 1620 |
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|     | <211> TYPE: |              |             |              |              |             |      |
|     | <213> ORGAN |              | icial Compo | 000          |              |             |      |
|     | <220> FEATU |              | tciai beque | ii Ce        |              |             |      |
|     |             |              | M. Saguence | e of a syntl | netic lucife | erase       |      |
|     | <400> SEQUE |              | M. Dequence | s or a sync. | iccic iucii  |             |      |
| 127 | 24002 SEQUI | accasasasas  | aataatataa  | ggcccagagc   | ctctgcaccc   | attogaagac  | 60   |
| 130 | atgatgaaac  | gtgaaaagaa   | attacataat  | ctgcgtaaac   | attotoactt   | acctcaaacc  | 120  |
| 120 | ctgategetg  | togtgagargra | greecgreet  | tcttataagg   | aatttttcca   | agetactate  | 180  |
| 139 | atattagasa  | natototosa   | taattaaaat  | tacaaaatga   | acceptatant  | carcatttat  | 240  |
| 140 | cegueggeee  | aatttttt     | tttcatcaca  | gtgattgccg   | attactacat   | cagcataatt  | 300  |
| 141 | guigagaata  | tanatanata   | ttatatooca  | gacgagttgt   | acaaaatcat   | aggtattage  | 360  |
|     |             |              |             | attctgaata   |              |             | 420  |
|     |             |              |             |              |              |             | 480  |
| 144 | egtactaact  | teateaageg   | Cattateatt  | ctggataccg   | atatogataa   | attagagagat | 540  |
| 145 | gagagettge  | caaactttat   | ttetegitat  | agcgacggta   | acategecaa   | citcaagcci  | 600  |
|     |             |              |             | gctattttgt   |              |             | 660  |
| 14/ | ctgcctaaag  | gcgtgatgca   | gactcaccaa  | aatatctgtg   | teegettgat   | cttaggeettg | 720  |
| 148 | gacccacgtg  | tgggtactca   | gttgatecet  | ggcgtgactg   | teetggtgta   | citycoatto  | 780  |
| 149 | tttcacgcct  | teggtttttc   | tattaccctg  | ggctatttca   | tggtcggttt   | gegegegate  | 840  |
| 150 | atgittcgtc  | gcttcgatca   | agaagccttt  | ctgaaggcca   | ttcaagacta   | egaggteegt  | 900  |
| 151 | agcgtgatca  | acgtcccttc   | tgtgattttg  | ttcctgagca   | aateteeatt   | ggicgalaag  | 960  |
| 152 | tatgacctga  | gcagcttgcg   | cgaactgtgc  | tgtggcgctg   | cccctttggc   | taaagaggty  | 1020 |
|     |             |              |             | ccaggtatcc   |              |             |      |
| 154 | gagagcacct  | ctgctaacat   | tcatagcttg  | cgtgatgagt   | tcaaatctgg   | cagectgggt  | 1080 |
| 155 | cgcgtgactc  | ctttgatggc   | cgctaagatc  | gccgaccgtg   | agaccggcaa   | agctctgggt  | 1140 |
| 156 | ccaaatcaag  | tcggcgaatt   | gtgtattaag  | ggtcctatgg   | tgtctaaagg   | ctacgtcaac  | 1200 |
| 157 | aatgtggagg  | ccactaagga   | agctattgat  | gacgatggtt   | ggctgcacag   | cggcgacttt  | 1260 |
| 158 | ggttattacg  | atgaggacga   | acatttctat  | gtcgtcgatc   | gctacaaaga   | gttgattaag  | 1320 |
| 159 | tataaaggct  | ctcaagtcgc   | cccagctgag  | ctggaagaaa   | tcttgctgaa   | gaacccttgc  | 1380 |
| 160 | attegtgacg  | tggccgtcgt   | gggtatccca  | gatttggaag   | ctggcgagct   | gcctagcgcc  | 1440 |
| 161 | tttgtcgtga  | aacaaccagg   | caaggaaatt  | accgctaaag   | aggtctacga   | ctatttggcc  | 1500 |
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| 163<br>164            | gagegegtgt eteacactaa gt<br>egeaatgtea eeggeaaaat ta<br>ggtgge  |            |            |            |            | 1560<br>1620<br>1626 |  |  |  |
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|                       | 172 <223> OTHER INFORMATION: Sequence of a synthetic luciferase |            |            |            |            |                      |  |  |  |
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|                       | ctcaccgctg gtgagatgct gt<br>ctcgtggacg tcgtgggtga cg            |            |            |            |            | 180                  |  |  |  |
|                       | ctqttqqccc aaaqcctqca ta  |            |            |            |            | 240                  |  |  |  |
|                       | qctqagaata acactcgctt tt  |            | -          |            |            | 300                  |  |  |  |
|                       | gtcgcccctg tgaatgaatc tt  |            |            |            |            | 360                  |  |  |  |
|                       | aaacctcaaa tcgtctttac ta  |            |            |            |            | 420                  |  |  |  |
|                       | cgtactaatt tcatcaaacg ca  |            |            |            |            | 480                  |  |  |  |
|                       | gagagettge ctaactttat ct  |            |            |            |            | 540                  |  |  |  |
|                       | ctgcattttq atccagtcga qc  |            |            | -          | -          | 600                  |  |  |  |
|                       | ttqcctaaaq qtqtcatqca qa  |            |            |            |            | 660                  |  |  |  |
|                       | qacceteqtq tqqqtactca at  |            |            |            |            | 720                  |  |  |  |
|                       | tttcacqcct ttqgtttttc ta  |            |            |            |            | 780                  |  |  |  |
|                       | atgtttcgtc gcttcgacca ag  |            |            |            |            | 840                  |  |  |  |
|                       | totgtgatca atgtoccato to  | , ,        |            |            |            | 900                  |  |  |  |
|                       | tatgatetga geagettgeg tg  |            |            |            |            | 960                  |  |  |  |
|                       | googaggtog otgotaagog to  |            |            |            |            | 1020                 |  |  |  |
|                       | gagagcactt ctgccaacat co  |            |            |            |            | 1080                 |  |  |  |
|                       | egegtgacee etttgatgge tg  |            |            |            |            | 1140                 |  |  |  |
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|                       | aatgtggagg ccactaaaga ag  |            |            |            |            | 1260                 |  |  |  |
| 196                   | ggttattatg atgaggacga ac  | cacttctat  | gtggtcgatc | gctataaaga | attgattaag | 1320                 |  |  |  |
| 197                   | tacaaagget etcaagtege co  | cagctgaa   | ctggaagaaa | ttttgctgaa | gaacccttgt | 1380                 |  |  |  |
| 198                   | attegegacg tggeegtegt gg  | gtatecca   | gacttggaag | ctggcgagtt | gectagegee | 1440                 |  |  |  |
| 199                   | tttgtggtga aacaacctgg ca  | aggagatt   | actgctaagg | aggtctacga | ctatttggcc | 1500                 |  |  |  |
| 200                   | gagcgcgtgt ctcacactaa at  | atctgcgt   | ggcggcgtcc | gcttcgtcga | ttctatccct | 1560                 |  |  |  |
| 201                   | cgcaacgtca ccggcaagat ca  | ictcgtaaa  | gagttgctga | aacaattgct | cgaaaaagct | 1620                 |  |  |  |
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|                       | <212> TYPE: DNA   |            |            |            |            |                      |  |  |  |
|                       | <213> ORGANISM: Artifici  | .al Sequen | ice        |            |            |                      |  |  |  |
|                       | 209 <220> FEATURE:  |            |            |            |            |                      |  |  |  |
|                       | 210 <223> OTHER INFORMATION: Sequence of a synthetic luciferase |            |            |            |            |                      |  |  |  |
| 212 <400> SEQUENCE: 6 |   |            |            |            |            |                      |  |  |  |
|                       | atgatgaaac gcgaaaagaa cg  |            |            |            |            | 60                   |  |  |  |
|                       | ctcaccgctg gtgagatgct ct  |            |            |            |            | 120                  |  |  |  |
|                       | ctcgtggacg tcgtgggaga cg  |            |            |            |            | 180                  |  |  |  |
| 210                   | ctgttggccc aaagcctcca ta  | lattgtgga  | tacaaaatga | acgatgtggt | gagcatttgt | 240                  |  |  |  |
|                       |   |            |            |            |            |                      |  |  |  |

PATENT APPLICATION: US/09/645,706 TIME: 12:02:11 Input Set : A:\341-005US1.txt.txt Output Set: N:\CRF3\09062000\1645706.raw 300 217 gctgagaata acactcgctt ctttatccct gttatcgctg cttggtacat cggcatgatt 360 218 gtcgcccctg tgaatgaatc ttacatccca gatgagctgt gtaaggttat gggtattagc 420 219 aaacctcaaa tcgtctttac taccaaaaat atcctgaata aggtcttgga agtccagtct 220 cgtactaact tcatcaaacg catcattatt ctggataccg tcgaaaacat ccatggctgt 480 221 gagageetge ctaactteat etetegttae agegatggta atategetaa ttteaaacea 540 600 222 etgeattttg atccagtega geaagtggee getattttgt getetteegg eaceaetggt 223 ttgcctaaag gtgtcatgca gactcaccag aatatctgtg tgcgtttgat ccacgctctc 660 224 gaccetegtg tgggtactea attgatecet ggegtgaetg tgetggtgta tetgeettte 720 225 tttcacgcct ttggtttttc tattaccctg ggctatttca tggtcggctt gcgtgtcatc 780 226 atgtttegte gettegaeca agaageette ttgaaggeta tteaagaeta egaggtgegt 840 227 totgtoatca atgtocotto agtoattttg ttootgagea aatotoottt ggttgacaag 900 228 tatgatotga goagottgog tgagetgtgo tgtggogotg otcotttggo caaagaagtg 960 229 gccgaggtcg ctgctaagcg tctgaacctc cctggtatcc gctgcggttt tggtttgact 1020 230 gagagcactt ctgctaacat ccatagcttg cgagacgagt ttaagtctgg tagcctgggt 1080 231 egeqtqacte etettatqqe tqcaaagate geegacegtg agaceggeaa agcactggge 1140 232 ccaaatcaag toggtgaatt gtgtattaag ggccctatgg tototaaagg ctacgtgaac 1200 233 aatgtggagg ccactaaaga agccattgat gatgatggct ggctccatag cggcgacttc 1260 234 ggttactatg atgaggacga acacttctat gtggtcgatc gctacaaaga attgattaag 1320 235 tacaaagget etcaagtege eccageegaa etggaagaaa ttttgetgaa gaaceettgt 1380 236 atccgcgacg tggccgtcgt gggtatccca gacttggaag ctggtgagtt gcctagcgcc 1440 237 tttgtggtga aacaacctgg aaaggagatc actgctaagg aggtctacga ctatttggcc 1500 1560 238 gagogogtgt ctcacaccaa atatotgogt ggoggogtoc gottogtoga ttccatecca 1620 239 cgcaacgtga ccggtaagat cactcgtaaa gaattgctga agcaactcct cgaaaaagct 1626 240 gacaac 242 <210> SEQ ID NO: 7 243 <211> LENGTH: 1626 244 <212> TYPE: DNA 245 <213> ORGANISM: Artificial Sequence 247 <220> FEATURE: 248 <223> OTHER INFORMATION: Sequence of a synthetic luciferase 250 <400> SEQUENCE: 7 251 atgatgaaac gcgaaaagaa cgtgatctac ggcccagaac cactgcatcc actggaagac 252 ctcaccqctq qtqaqatqct cttccqaqca ctqcgtaaac atagtcacct ccctcaagca 120 253 ctcgtggacg tcgtgggaga cgagagcctc tcctacaaag aatttttcga agctactgtg 180 254 ctqttqqccc aaaqcctcca taattqtqqq tacaaaatqa acgatqtqgt gagcatttgt 240 255 getgagaata acactegett etttatteet gtaategetg ettggtacat eggeatgatt 300 256 gtcgcccctg tgaatgaatc ttacatccca gatgagetgt gtaaggttat gggtattagc 257 aaacctcaaa tcgtctttac taccaaaaac atcttgaata aggtcttgga agtccagtct 420 258 cgtactaact tcatcaaacg catcattatt ctggataccg tcgaaaacat ccacggctgt 259 gagageetee etaaetteat etetegttae agegatggta atategetaa titteaageee 540 260 ttgcattttg atccagtcga gcaagtggcc gctattttgt gctcctccgg caccactggt 261 ttgcctaaag gtgtcatgca gactcaccag aatatctgtg tgcgtttgat ccacgctctc 262 gaccctcgtg tgggtactca attgatccct ggcgtgactg tgctggtgta tctgcctttc 660 263 tttcacgect ttggtttete tattacectg ggetatttea tggteggett gegtgteate 264 atgtttegte gettegacea agaageette ttgaaggeta tteaagacta egaggtgegt 780 265 teegtgatea aegteeette agteattttg tteetgagea aateteettt ggttgacaag 900 266 tatgatotga goagottgog tgagotgtgo tgtggogotg otootttggo caaagaagtg 960 1020 267 gccgaggtcg ctgctaagcg tctgaacctc cctggtatcc gctgcggttt tggtttgact 268 gagagcactt ctgctaacat ccatagcttg cgagacgagt ttaagtctgg tagcctgggt

RAW SEQUENCE LISTING

ZF.Y.1.

DATE: 09/06/2000

## <u>Please Note:</u>

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

. . .

DATE: 09/06/2000 TIME: 12:02:12

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/645,706

Input Set : A:\341-005US1.txt.txt Output Set: N:\CRF3\09062000\1645706.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:4423 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:228 L:4581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:242 L:4618 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:245

- 11111666666